

Amendments to the Specification:

Please amend the Specification as follows:

Please amend all references to “Figure 9” to -- Figures 9A &9B -- as follows

At page 14 of the specification, use these replacement paragraphs, in accordance with 37 CFR 1.121(b).

FIGURES 9A & 9B is ~~a~~re black and white photographs of laminin digested with elastase, separated by SDS-PAGE and following interaction with biotinylated A $\beta$  (1-40). A ~55 kilodalton laminin fragment (arrow) that binds biotinylated A $\beta$  was identified and sequenced. Note also the presence of a ~130 kDa fragment (arrowheads) that binds A $\beta$  following 1.5 hours of elastase digestion (lane 2). Panel AFigure 9A is a ligand blot using biotinylated A $\beta$  as a probe, whereas panel BFigure 9B is Coomassie blue staining of the same blot in Panel AFigure 9A to locate the specific band(s) for sequencing.

FIGURE 10 shows the complete amino acid sequence of the mouse laminin A chain. Sequencing of the ~55 kilodalton A $\beta$ -binding band shown in Figure 9Figures 9A & 9B leads to the identification of an 11 amino acid segment (underline and arrowhead) within the laminin A chain. This A $\beta$  binding region of laminin is situated within the globular domain repeats of the laminin A chain.

At page 41 of the specification (Example 6), use these replacement paragraphs, in accordance with 37 CFR 1.121(b).

In Figure 9, Panel AFigure 9A represents an A $\beta$  ligand blot whereas panel BFigure 9B represents the equivalent Coomassie blue stained blot. As shown in Figure 9, Panel AFigure 9A (lanes 2 and 3), elastase-digested laminin produced multiple protein fragments which bound biotinylated A $\beta$  (1-40). Panel AFigure 9A, lane 1 represents undigested mouse EHS laminin, whereas lanes 2 and 3 represents laminin which had been digested with elastase for 1.5 hours or

2.5 hours, respectively. Panel AFigure 9A, lane 4 represents elastase digestion for 2.5 hours in the absence of laminin. Undigested laminin (Fig. 9, Panel AFigure 9A, lane 1) which interacted with A $\beta$  included multiple bands from  $> \sim 400$  kDa to  $> \sim 86$  kDa, with the most prominent A $\beta$ -interaction occurring with intact laminin (i.e.  $\sim 400$  kDa). Elastase-resistant laminin protein fragments which interacted with A $\beta$  (Fig. 9, Panel AFigure 9A, lanes 2 and 3) included fragments of  $> \sim 400$  kDa,  $\sim 130$  kDa (arrowhead),  $\sim 80$ - $90$  kDa,  $\sim 65$  kDa and a prominent band at  $\sim 55$  kDa (arrow). The interaction of these elastase-resistant laminin protein fragments with A $\beta$  were only observed under non-reducing conditions suggesting that the A $\beta$  interaction was also conformation dependent. The 130kDa elastase resistant laminin fragment which interacts with A $\beta$ , is also believed to be part of the E8 fragment (see Figure 11), and is the same protein fragment of laminin that appears to be present in human serum and cerebrospinal fluid (see Examples 10 and 11). Figure 9, Panel AFigure 9A, lane 4 demonstrates that the band observed at  $\sim 29$  kDa represents non-specific A $\beta$  binding due to the presence of the elastase enzyme alone.

Figure 9, Panel BFigure 9B demonstrates all of the multiple protein bands which were stained by Coomassie blue. Note, for example, in Panel BFigure 9B, lanes 2 and 3, that elastase digestion of laminin produced multiple protein fragments between  $\sim 55$  kDa and  $\sim 90$  kDa which did not bind A $\beta$ , and were not observed in the A $\beta$  ligand blot (Fig. 9, Panel AFigure 9A, lanes 2 and 3).

Please amend the Drawings as follows:

Figure 9 has been re-numbered to appear as Figures 9A and 9B (see attached proposed drawing correction sheet).